

Joint Analysis of Longitudinal and Survival AIDS Data – a Bayesian perspective

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Spatial Joint Analysis for AIDS Data

• Longitudinal Model

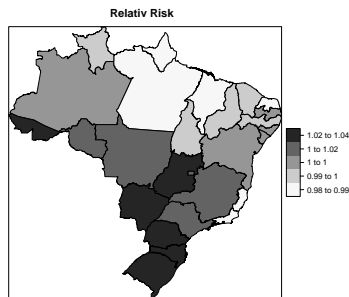
$$y_{ikj} = \mu_{ik}(t_{ikj}) + W_{1ik}(t_{ikj}) + \epsilon_{ikj}$$

- $y_{ikj} = \sqrt{\text{CD4}}$: j -th count for the individual $i = 1, \dots, n_k$ living in the $k = 1, \dots, K$ region
- $W_{ik}(t) = (W_{1ik}(t), W_{2ik}(t))$: zero-mean latent bivariate Gaussian process
- $W_{1ik} = U_{1ik} + U_{2ik}t_{ikj}$
- $(U_{1ik}, U_{2ik}) \sim N_2(\mathbf{0}, \Sigma)$

• Survival Model

$$T_{ik} \sim \text{Weibull}(r, \mu_{ik}(t))$$

- $\log(\mu_{ik}(t)) = \mathbf{x}_{2ik}^T(t)\beta_2 + W_{2ik}(t) + Q_k$
- $W_{2ik} = \gamma_1 U_{1ik} + \gamma_2 U_{2ik}$.
- $Q_k | \sigma_Q^2 \sim \text{ICAR}(\sigma_Q^2)$: spatial frailty



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